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Figure 1a

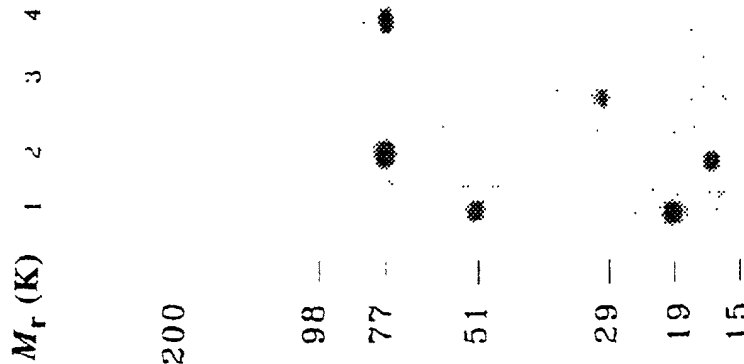
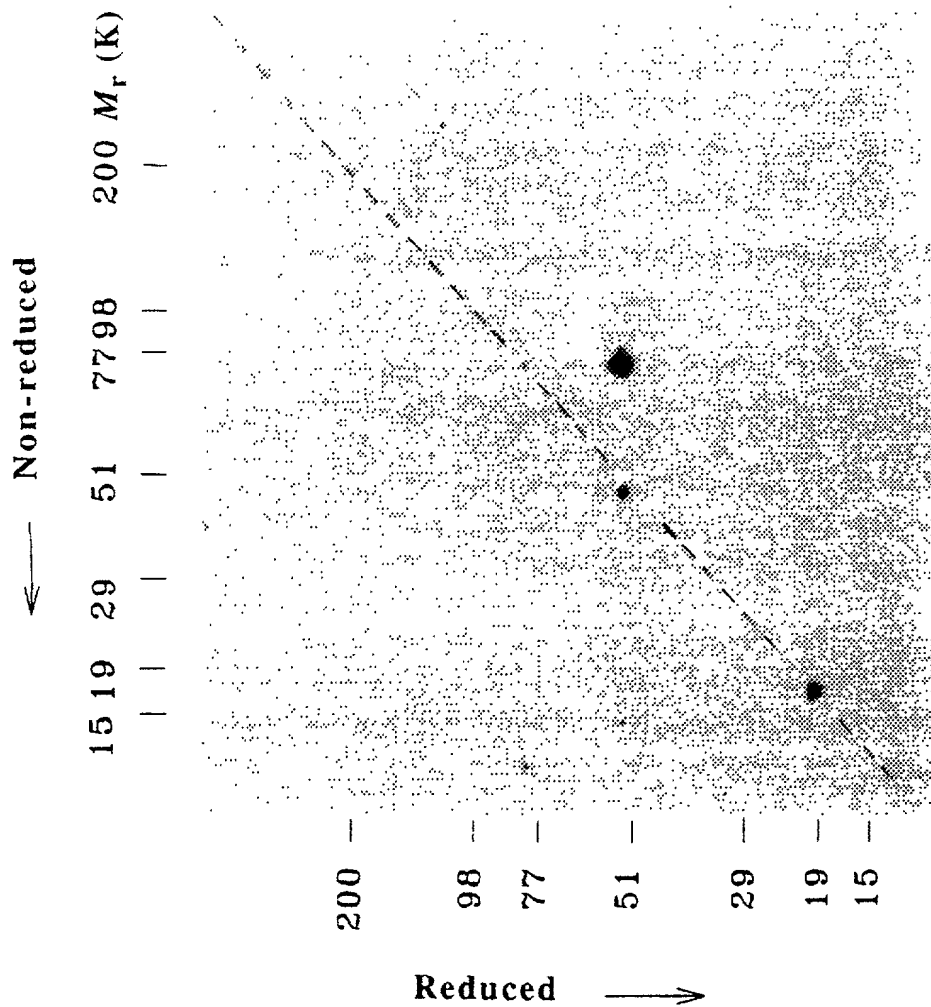


Figure 1b



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Figure 2

MASP-2		C1r/C1s →		TPLGPKWPEPVFGRLASPGFPGEYANDOEERRWLTAPPGYRLRLYFTHFDLELSHLCEYDFVKLSSGAKVLATLGGQESTDTERAPGKDT	90
MASP-1	HTVELNNMFGQIQSPGYPDSYPSDSEVTWNITVPDGFRIKLYFMHFNLESSYLCEYDYVKVETEDQVLATPGRETDDTEQTPGQEV				87
C1r	SIPIPQKLFGVEVTSPLFPKYPNNFETTTTITVPTGYRVKLVFQQFDLEPSEGCFYDYVKISADKKSLGRFCGQLGSPGNPPGKKE				87
C1s	EPTMYGEILSPNYPQAYPSEVEKSWDIEVPEGYGIHLYFTHLDIELSENAYDSVQIISGDTTEGRLEGQRSSNNPHSPIVEE				83
		* * * * *			
MASP-2		ECF →		FYSLGSSLDITFRSDYSNEKP FTGFEAFYAEDIDEQ VAPGEA PTCDHHCHNHLGGFY@SCRAGYVLHRNKRTGSALES	170
MASP-1	VLSPGFSMSITFRSDFSNEER FTGFDAYMAVDVDECK EREDEE LSCDHYCHNYIGGYE@SCRFYILHTDNRTERVECS				167
C1r	FMSQGNKMLLTFTDSDSNEENGITIMFYKGLAYYQAVDLDEASRSKSGEEDPQCCQHLCHNYVGGYF@SCRPYELQEDRHS@QAECS				177
C1s	FQVPYNKLQVIFKSDFSNEER FTGFAAYVATDINET DFVD VPCSHFCNNFIGGYF@SCPEYFLHDDMKNEGVNCS				161
		* * * * *			
MASP-2		- C1r/C1s →		GQVFTQSGELSSPEYPRPYPKLSSCTYSISLEEGFSVILDFV ESFDVET HPETLCPYDFLKIQTDRREEHGPF@GKTLPHR IETKS	256
MASP-1	DNLFTQRTGVITSPDPNPNPKSSE@LYTIELEEGFMVNLQFE DIFDIED HPEVP@PYDYIKIKVGPKVLGPF@GGEKAPEP ISTQS				253
C1r	SELYTEASGYISSLEYPRSYPPDLR@NYSIRVERGLTLHLKFL EPPDIDD HQQVHC@PYDQLQIYANGKNIGEF@GKQRPDP LDTSS				263
C1s	GDVFTALIGEIASPNYPKYPENSRC@EYQIRLEKGFQVVTLRREDFDVEAADSAGN@ LDSLVFVAGDRQFGPY@GHGFPGLNIETKS				250
		* * * * *			
MASP-2		CCP-1 →		NTVTITFVTDSEGDHTGKWIHYTSTAQ@PYPMAPPN GHVSPVQAKYILKDSFSIF@ETGYELLQGHLPKLSFTAV@QKDGSDWRPMPA	345
MASP-1	HSVLILFHDNSNGENRGWRLSYRAAGNE@PELQPPVH GKIEPSQAKYFFKQVLVS@DTGYKVLKDNVEMDT@QIE@LKDGTWSNKIPT				342
C1r	NAVDLLFFTDESGLSRGWKLRYTTEI@K@PQKTLDEFTII@NLQ@PYQ@FRDYFIAT@KQGYQLIEGNOVLHSFTAV@QDDGTWHRAMPR				353
C1s	NALDII@TDLTGQKKGWKLRYHGDPM@P@KEDTPN SVWEPAKAKYVFRD@VQIT@LDGFV@V@GRV@G@TSFYST@Q@SNGKWSNSK@L@K				338
		* * * * *			
MASP-2		CCP-2 →		ESIVD@CGPPDDLPSGRVEYITGPGVTTYKAVIQYSC@ETFYTM KVNDGKYV@EADGFWTSSKGEKSLPV@EPV@GLS ARTT	426
MASP-1	CKIVD@RAPGELEHGLITFSTRNNLT@YKSEIKYS@Q@EPYKML NNNTGIYT@SAQGV@M@NKV@LGRSLPT@L@PV@GLPKFSRKL				426
C1r	CKIKD@Q@P@RN@L@P@NGD@FRYTT@TM@VNTYKARIQY@HEPYK@M@TRAGSRESE@Q@VYT@TAQGIW@KNE@Q@GEKIP@L@PV@G@K@P@V@P@VEQ				443
C1s	C@PV@D@G@I@P@ESIENGKVE DPESTLFGSVIRYTT@E@PY@YME NGGGGEYH@CAGNSW@VNEVL@G@PEL@P@K@V@P@V@G@V@P@P@F@EE				419
		* * * * *			
MASP-2		serine protease →		GGRIYGGQKAKPGDFPWQVLILGGTTA AGALLYDNWVLTAAH AVYEQKH@DASALDIRM@TLKRLSPHYTQAWSEAVFIHEG	507
MASP-1	MARIFNGRPAQKGTT@PWIA@MLSHLNG@PFCGGSLLGSSWIVTA@AHCLH@SLD@PKD@PTL@RDS@LL@SPSD FKII@LGK@H@W@L@R@SD@ENE@Q@HLG				515
C1r	RQRIIGGQKAKMGNFPWQVFTNIHGRG GGALLGDRWILTAAH TLYPKEH@EAQ@NASL@DVFLGHT@NVEEL@MKL@GNHP IRRV				523
C1s	KQRIIGGSDADIKNFPWQVFPDNPWA GGALINEY@WVLTAAH VVEGNREPTMYVGSTSVQTSRLAKSKMLT PEHVFIHPG				498
		* * * * *			
MASP-2		YTHDAG		FDNDIALIKL@NKV@VINSNIT@P@L@P@RKE@ESFMRTDDIGTASG@WGLT@RGFLARNL@MYVDI@PVDH@K@C@TAA@YEK	589
MASP-1	VKHTTLHPKYDPNTFENDVALVELLES@PVLNAF@VMP@ILP EGPQ@EGAMVIVSGW@KQFL@R@P@PETLMEIEI@PVDHST@Q@KAY				599
C1r	SVHPDYRQDESYN FEGDIALLE@L@NSVTL@G@NLL@P@ILP DNDTFYDLGLM@GVVSGFGVME@EK I@HDLRFVRL@P@VAN@P@Q@E@N WLR				608
C1s	WKLLEV PEGRTN FDNDIALVRLKDPVKMGPTVSP@ILPGTSSDY@NLM@DGL@LISG@WGRTEKRDRAVRLK@ARLP@VAPL@R@K@KEV@KVE				586
		* * * * *			
MASP-2		PPYPRG		SVTANMLC@AGLES@G@K@D@S@RGD@SG@ALV@FLDS ETERW@FVG@IVSWG@SMN@G@EAG@QYGVYTKVINI@P@WIENIISDF	671
MASP-1	APLKK K@TRDMIC@AGEK@G@K@D@C@SGD@SG@PMVTLNR ERGQWYLVGT@VSWG@ D@G@K@DRYGVYSYI@H@NK@W@I@Q@RTV@GV@RN				680
C1r	GKNRMD VFSQNMFC@AGH@PSL@KQD@C@Q@GDS@GVFAVRDP N@DRW@VAT@GIVSWG@ G@SRG YG@FYTKV@LNYVDWIK@KEME@EED				688
C1s	KPTADAEAYVFTPNM@C@AGGEK G@MDS@K@GDS@G@AF@V@Q@PNDKTKFYA@AGLV@SWG@ Q@CGT YGLYTRV@KNYVDWIM@KTM@QENST@PRED				673
		* * * * *			

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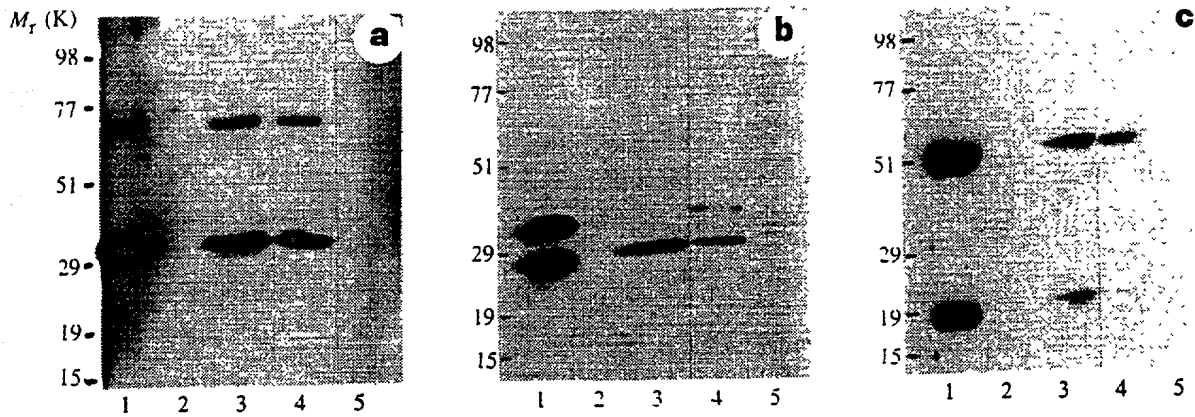
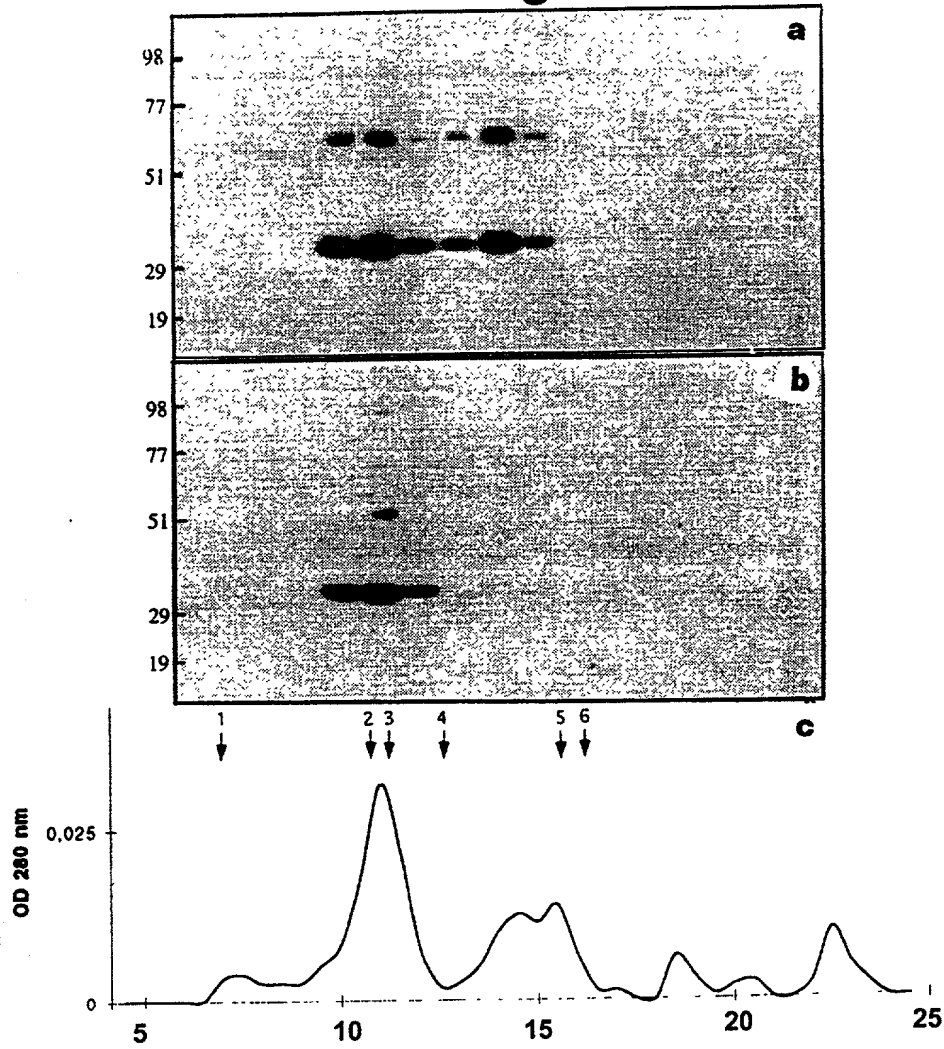


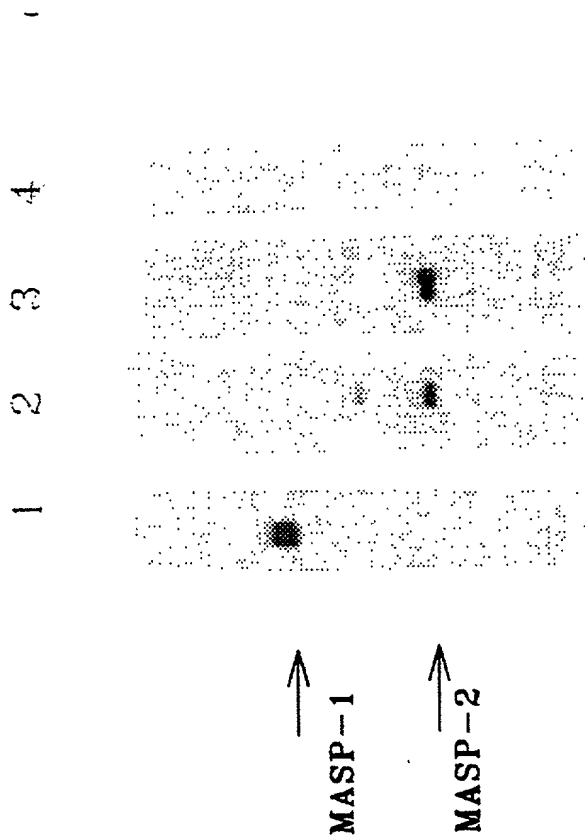
Figure 3b



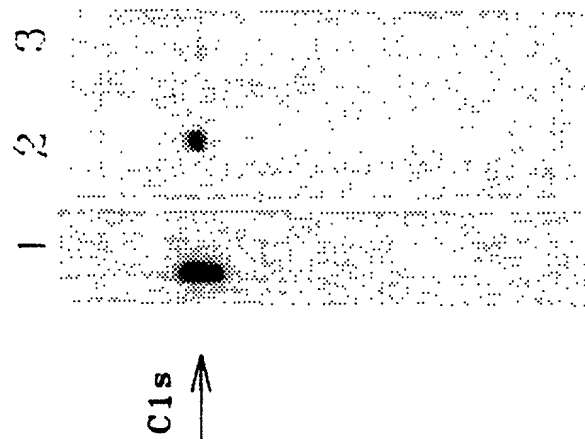
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Figure 4

Blot of MBL preparation

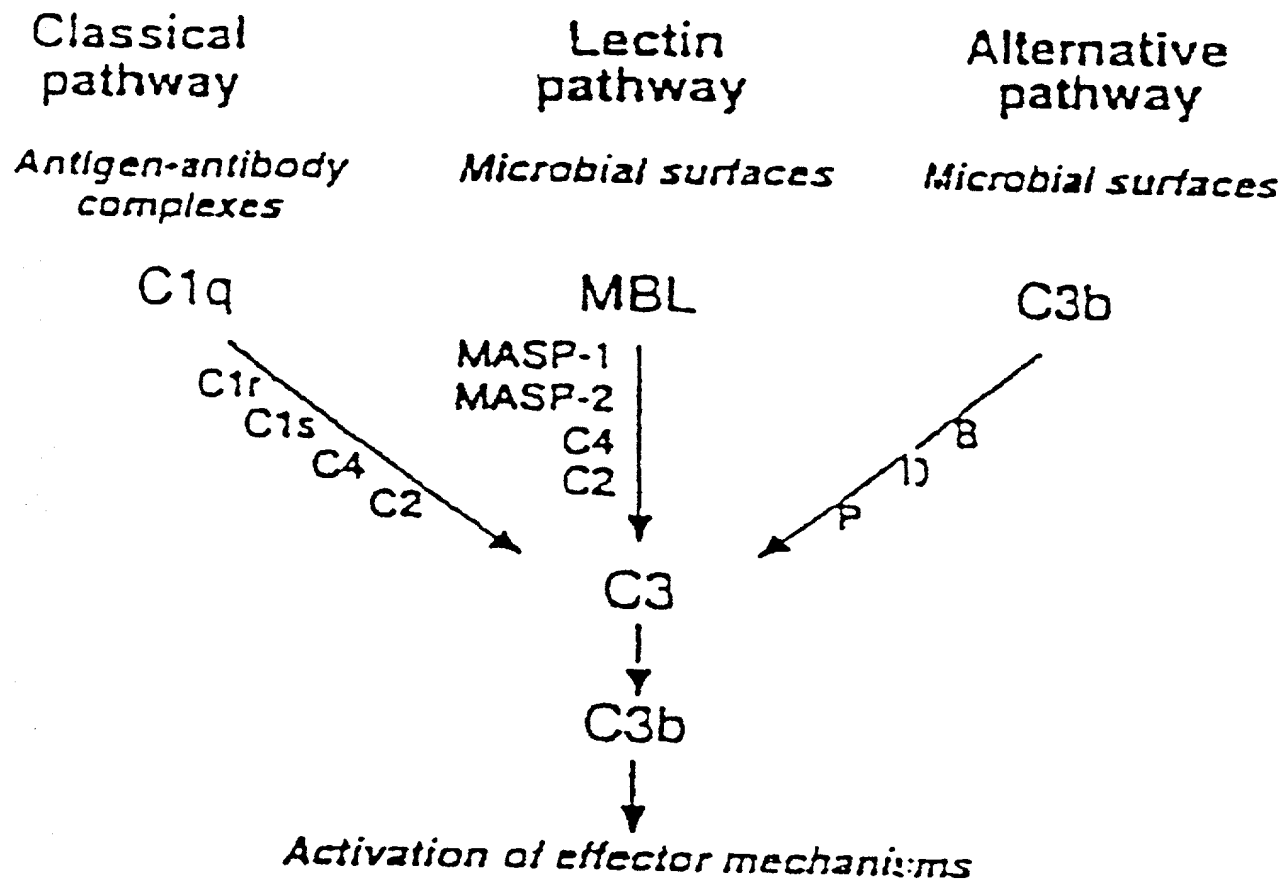


Blot of C1



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Figure 5



[illegible]